/FW16

# CRF Errors Edited by the STIC Systems Branch

	Number: 09/870,375A	CRF Edit Date: 9/20/04 Edited by: AZ
	Realigned nuclear acid/amino cd_numbers/text text "syrapped" to the next line	in cases where the sequence
	Corrected the SEQ ID NO. Sequence numbers of	edited were:
	Inserted or corrected a nucleic number at the en	d of a nucleic line. SEQ ID
<del></del>	Deleted: invalid beginning/end-of-file text;	page numbers
	Inserted mandatory headings/numeric identifier	rs, specifically:
	Moved responses to same line as heading/numer	ic identifier, specifically:
<u>J</u>	Others Segvene 22- charged (4007	X8 22107



IFW16

RAW SEQUENCE LISTING

DATE: 09/20/2004

PATENT APPLICATION: US/09/870,375A

TIME: 11:52:50

Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\09202004\1870375A.raw

- 3 <110> APPLICANT: Greuber et al.
- 5 <120> TITLE OF INVENTION: CHIMERIC PLANT PROMOTERS AND PLANTS CONTAINING THE SAME
- 7 <130> FILE REFERENCE: 18433/2022
- 9 <140 > CURRENT APPLICATION NUMBER: 09/870,375A
- 10 <141> CURRENT FILING DATE: 2001-05-30
- 12 <150> PRIOR APPLICATION NUMBER: PCT/IB00/01383
- 13 <151> PRIOR FILING DATE: 2000-09-28
- 15 <160> NUMBER OF SEQ ID NOS: 38
- 17 <170> SOFTWARE: PatentIn Ver. 2.1
- 19 <210> SEQ ID NO: 1
- 20 <211> LENGTH: 417
- 21 <212> TYPE: DNA
- 22 <213> ORGANISM: Triticum aestivum
- 24 <220> FEATURE:
- 25 <221> NAME/KEY: misc feature
- 26 <222> LOCATION: (22)..(29)
- 27 <223> OTHER INFORMATION: Prolamine- like box
- 29 <220> FEATURE:
- 30 <221> NAME/KEY: misc feature
- 31 <222> LOCATION: (70)..(73)
- 32 <223> OTHER INFORMATION: GATA box
- 34 <220> FEATURE:
- 35 <221> NAME/KEY: misc\_feature
- 36 <222> LOCATION: (87)..(90)
- 37 <223> OTHER INFORMATION: GATA box
- 39 <220> FEATURE:
- 40 <221> NAME/KEY: misc feature
- 41 <222> LOCATION: (127)..(133)
- 42 <223 > OTHER INFORMATION: Prolamine-like box
- 44 <220> FEATURE:
- 45 <221> NAME/KEY: misc\_feature
- 46 <222> LOCATION: (161)..(168)
- 47 <223> OTHER INFORMATION: G-like box
- 49 <220> FEATURE:
- 50 <221> NAME/KEY: enhancer
- 51 <222> LOCATION: (193)..(230)
- 52 <223> OTHER INFORMATION: Enhancer box
- 54 <220> FEATURE:
- 55 <221> NAME/KEY: TATA signal
- 56 <222> LOCATION: (349)..(355)
- 57 <223> OTHER INFORMATION: TATA box
- 59 <220> FEATURE:
- 60 <221> NAME/KEY: misc feature

PATENT APPLICATION: US/09/870,375A

DATE: 09/20/2004 TIME: 11:52:50

Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\09202004\1870375A.raw

- 61 <222> LOCATION: (379)
- 62 <223> OTHER INFORMATION: Transcription initiation site
- 64 <400> SEQUENCE: 1
- 65 agetttgagt ggeegtagat ttgcaaaage aatggetaae agacacatat tetgecaaae 60
- 66 cccaagaagg ataatcactt ttcttagata aaaaagaaca gaccaatata caaacatcca 120
- 67 cacttetgea aacaatacat cagaactagg attacgeega ttacgtgget ttageagact 180
- 68 gtccaaaaat ctgttttgca aagctccaat tgctccttgc ttatccagct tcttttgtgt 240
- 69 tggcaaactg cgcttttcca accgattttg ttcttctcgc gctttcttct taggctaaac 300
- 70 aaaceteace gtgeaegeag ceatggteet gaacetteae etegteeeta taaaageeta 360
- 71 gccaacette acaatettat cateacecae aacacegage accacaaact agagate 417
- 74 <210> SEQ ID NO: 2
- 75 <211> LENGTH: 181
- 76 <212> TYPE: DNA
- 77 <213> ORGANISM: Artificial Sequence
- 79 <220> FEATURE:
- 80 <223> OTHER INFORMATION: Description of Artificial Sequence: MPr1126
- 81 promoter
- 83 <220> FEATURE:
- 84 <221> NAME/KEY: TATA signal
- 85 <222> LOCATION: (113)..(119)
- 86 <223> OTHER INFORMATION: TATA box
- 88 <220> FEATURE:
- 89 <221> NAME/KEY: misc feature
- 90 <222> LOCATION: (143)
- 91 <223> OTHER INFORMATION: Transcription Initiation site
- 93 <400> SEQUENCE: 2
- 94 gtgttggcaa actgcgcttt tccaaccgat tttgttcttc tcgcgctttc ttcttaggct 60
- 95 aaacaaacct caccgtgcac gcagccatgg teetgaacct teacetegte cetataaaag 120
- 96 cctagccaac cttcacaatc ttatcatcac ccacaacacc gagcaccaca aactagagat 180
- 97 c
- 100 <210> SEQ ID NO: 3
- 101 <211> LENGTH: 244
- 102 <212> TYPE: DNA
- 103 <213> ORGANISM: Artificial Sequence
- 105 <220> FEATURE:
- 106 <223> OTHER INFORMATION: Description of Artificial Sequence: MPr1127
- 107 promoter
- 109 <220> FEATURE:
- 110 <221> NAME/KEY: enhancer
- 111 <222> LOCATION: (20)..(57)
- 112 <223> OTHER INFORMATION: Enhancer box
- 114 <220> FEATURE:
- 115 <221> NAME/KEY: TATA signal
- 116 <222> LOCATION: (176)..(182)
- 117 <223> OTHER INFORMATION: TATA box
- 119 <220> FEATURE:
- 120 <221> NAME/KEY: misc feature
- 121 <222> LOCATION: (206)
- 122 <223> OTHER INFORMATION: Transcription Initiation site

DATE: 09/20/2004 TIME: 11:52:50

PATENT APPLICATION: US/09/870,375A

Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\09202004\I870375A.raw

- 124 <400> SEQUENCE: 3
- 125 geagactgte caaaaatetg ttttgeaaag etceaattge teettgetta teeagettet 60
- 126 tttgtgttgg caaactgcgc ttttccaacc gattttgttc ttctcgcgct ttcttcttag 120
- 127 gctaaacaaa cctcaccgtg cacgcagcca tggtcctgaa ccttcacctc gtccctataa 180
- 128 aagectagee aacetteaca atettateat cacceacaac acegageace acaaactaga 240
- 129 gatc
- 132 <210> SEQ ID NO: 4
- 133 <211> LENGTH: 277 134 <212> TYPE: DNA
- 135 <213> ORGANISM: Artificial Sequence
- 137 <220> FEATURE:
- 138 <223> OTHER INFORMATION: Description of Artificial Sequence:MPr1128
- 139 promoter
- 141 <220> FEATURE:
- 142 <221> NAME/KEY: misc feature
- 143 <222> LOCATION: (21)..(28)
- 144 <223> OTHER INFORMATION: G-like box
- 146 <220> FEATURE:
- 147 <221> NAME/KEY: enhancer
- 148 <222> LOCATION: (53)..(90)
- 149 <223> OTHER INFORMATION: Enhancer box
- 151 <220> FEATURE:
- 152 <221> NAME/KEY: TATA signal
- 153 <222> LOCATION: (209)..(215)
- 154 <223> OTHER INFORMATION: TATA box
- 156 <220> FEATURE:
- 157 <221> NAME/KEY: misc feature
- 158 <222> LOCATION: (239)
- 159 <223> OTHER INFORMATION: Transcription Initiation Site
- 161 <400> SEQUENCE: 4
- 162 cagaactagg attacgccga ttacgtggct ttagcagact gtccaaaaaat ctgttttgca 60
- 163 aageteeaat tgeteettge ttateeaget tettttgtgt tggeaaaetg egetttteea 120
- 164 accepatiting ticticized gettictict taggetaaac aaaceteace gigeaegeag 180
- 165 ccatggtect gaacetteae etegteeeta taaaageeta geeaaeette acaatettat 240
- 166 catcacccac aacaccqaqc accacaaact agagatc 277
- 169 <210> SEQ ID NO: 5
- 170 <211> LENGTH: 472
- 171 <212> TYPE: DNA
- 172 <213> ORGANISM: Artificial Sequence
- 174 <220> FEATURE:
- 175 <223> OTHER INFORMATION: Description of Artificial Sequence: MPr1130
- promoter 176
- 178 <220> FEATURE:
- 179 <221> NAME/KEY: misc feature
- 180 <222> LOCATION: (22)..(29)
- 181 <223> OTHER INFORMATION: Prolamine-like box
- 183 <220> FEATURE:
- 184 <221> NAME/KEY: misc feature
- 185 <222> LOCATION: (70)..(73)

## RAW SEQUENCE LISTING DATE: 09/20/2004 PATENT APPLICATION: US/09/870,375A TIME: 11:52:50

Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\09202004\1870375A.raw

186 <223> OTHER INFORMATION: GATA box 188 <220> FEATURE: 189 <221> NAME/KEY: misc feature 190 <222> LOCATION: (87)..(90) 191 <223> OTHER INFORMATION: GATA box 193 <220> FEATURE: 194 <221> NAME/KEY: misc feature 195 <222> LOCATION: (127)..(133) 196 <223> OTHER INFORMATION: Prolamine-like box 198 <220> FEATURE: 199 <221> NAME/KEY: misc feature 200 <222> LOCATION: (161)..(168) 201 <223> OTHER INFORMATION: G-like box 203 <220> FEATURE: 204 <221> NAME/KEY: enhancer 205 <222> LOCATION: (193)..(230) 206 <223> OTHER INFORMATION: Enhancer box 208 <220> FEATURE: 209 <221> NAME/KEY: misc feature 210 <222> LOCATION: (314)..(368) 211 <223> OTHER INFORMATION: As2/As2/As1 box 213 <220> FEATURE: 214 <221> NAME/KEY: TATA signal 215 <222> LOCATION: (404)..(410) 216 <223> OTHER INFORMATION: TATA box 218 <220> FEATURE: 219 <221> NAME/KEY: misc feature 220 <222> LOCATION: (434) 221 <223> OTHER INFORMATION: Transcription Initiation Site 223 <400> SEQUENCE: 5 224 agetttgagt ggeegtagat ttgcaaaage aatggetaac agacacatat tetgecaaac 60 225 cccaagaagg ataatcactt ttcttagata aaaaagaaca gaccaatata caaacatcca 120 226 cacttetgea aacaatacat cagaactagg attacgeega ttacgtgget ttagcagaet 180 227 gtccaaaaat ctgttttgca aagctccaat tgctccttgc ttatccagct tcttttgtgt 240 228 tggcaaactg cgcttttcca accgattttg ttcttctcgc gctttcttct taggctaaac 300 229 aaacctcacc qtqattqatq tqatatcaaq attqatqtqa tatctccact qacqtaaqqq 360 230 atqacqcaca cqcaqccatq qtcctqaacc ttcacctcqt ccctataaaa qcctaqccaa 420 231 ccttcacaat cttatcatca cccacaacac cgagcaccac aaactagaga tc 234 <210> SEQ ID NO: 6 235 <211> LENGTH: 455 236 <212> TYPE: DNA 237 <213> ORGANISM: Artificial Sequence 239 <220> FEATURE: 240 <223> OTHER INFORMATION: Description of Artificial Sequence:MPr1131 promoter 243 <220> FEATURE: 244 <221> NAME/KEY: misc feature

245 <222> LOCATION: (22)..(29)

246 <223> OTHER INFORMATION: Prolamine-like box

### RAW SEQUENCE LISTING DATE: 09/20/2004 PATENT APPLICATION: US/09/870,375A TIME: 11:52:50

Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\09202004\I870375A.raw

```
248 <220> FEATURE:
249 <221> NAME/KEY: misc feature
250 <222> LOCATION: (70)..(73)
251 <223> OTHER INFORMATION: GATA box
253 <220> FEATURE:
254 <221> NAME/KEY: misc feature
255 <222> LOCATION: (87)..(90)
256 <223> OTHER INFORMATION: GATA box
258 <220> FEATURE:
259 <221> NAME/KEY: misc feature
260 <222> LOCATION: (127)..(133)
261 <223> OTHER INFORMATION: Prolamine-like box
263 <220> FEATURE:
264 <221> NAME/KEY: misc feature
265 <222> LOCATION: (161)..(168)
266 <223> OTHER INFORMATION: G-like box
270 <400> SEQUENCE: 6
271 agetttgagt ggeegtagat ttgcaaaage aatggetaac agacacatat tetgccaaac 60
272 cccaagaagg ataatcactt ttcttagata aaaaagaaca gaccaatata caaacatcca 120
273 cacttetgea aacaatacat cagaactagg attacgeega ttacgtgget ttageagaet 180
274 gtccaaaaat ctgttttgca aagctccaat tgctccttgc ttatccagct tcttttgtgt 240
275 tggcaaactg cgcttttcca accgattttg ttcttctcgc gctttcttct taggctaaac 300
276 aaacctcacc gtgattgatg tgatatctcc actgacgtaa gggatgacgc acacgcagcc 360
277 atggtcctga accttcacct cgtccctata aaagcctagc caaccttcac aatcttatca 420
                                                                       455
278 tcacccacaa caccgagcac cacaaactag agatc
280 <210> SEQ ID NO: 7
281 <211> LENGTH: 236
282 <212> TYPE: DNA
283 <213> ORGANISM: Artificial Sequence
285 <220> FEATURE:
286 <223> OTHER INFORMATION: Description of Artificial Sequence: MPr1133
287
          promoter
289 <220> FEATURE:
290 <221> NAME/KEY: misc feature
291 <222> LOCATION: (78)..(132)
292 <223> OTHER INFORMATION: As2/As2/As1 box
294 <220> FEATURE:
295 <221> NAME/KEY: misc feature
296 <222> LOCATION: (198)
297 <223> OTHER INFORMATION: Transcription Initiation Site
299 <400> SEQUENCE: 7
300 gtgttggcaa actgcgcttt tccaaccgat tttgttcttc tcgcgctttc ttcttaggct 60
301 aaacaaacct caccgtgatt gatgtgatat caagattgat gtgatatctc cactgacgta 120
302 agggatgacg cacacgcage catggteetg aacetteace tegteectat aaaagcetag 180
303 ccaacettea caatettate ateacecaca acacegagea ccacaaacta gagate
305 <210> SEQ ID NO: 8
306 <211> LENGTH: 299
```

308 <213> ORGANISM: Artificial Sequence

307 <212> TYPE: DNA

VERIFICATION SUMMARY

DATE: 09/20/2004

PATENT APPLICATION: US/09/870,375A

TIME: 11:52:51

Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\09202004\1870375A.raw

L:1100 M:112 C: (48) String data converted to lower case,



IFW16

RAW SEQUENCE LISTING

DATE: 09/17/2004

PATENT APPLICATION: US/09/870,375A

TIME: 09:55:40

**Does** Not Comply Corrected Diskette Needer

M12,4

Input Set : A:\final seq listing.TXT Output Set: N:\CRF4\09172004\1870375A.raw

```
3 <110> APPLICANT: Greuber et al.
```

- 5 <120> TITLE OF INVENTION: CHIMERIC PLANT PROMOTERS AND PLANTS CONTAINING THE SAME
- 7 <130> FILE REFERENCE: 18433/2022
- 9 <140> CURRENT APPLICATION NUMBER: 09/870,375A
- 10 <141> CURRENT FILING DATE: 2001-05-30
- 12 <150> PRIOR APPLICATION NUMBER: PCT/IB00/01383
- 13 <151> PRIOR FILING DATE: 2000-09-28
- 15 <160> NUMBER OF SEQ ID NOS: 38
- 17 <170> SOFTWARE: PatentIn Ver. 2.1

### ERRORED SEQUENCES

- 587 <210> SEQ ID NO: 16
- 588 <211> LENGTH: 96
- 589 <212> TYPE: DNA
- 590 <213> ORGANISM: Artificial Sequence
- 592 <220> FEATURE:
- 593 <223> OTHER INFORMATION: Description of Artificial Sequence:MPr1197
- promoter
- 596 <220> FEATURE:
- 597 <221> NAME/KEY: TATA signal
- 598 <222> LOCATION: (28)..(34)
- 599 <223> OTHER INFORMATION: TATA box
- 601 <220> FEATURE:
- 602 <221> NAME/KEY: misc feature
- 603 <222> LOCATION: (58)
- 604 <223> OTHER INFORMATION: Transcription Initiation Site
- 606 <400> SEQUENCE: 16
- 607 catggtcctg accettcace tegtccctat accettca caatcttate 60

#### E--> 608 atcacccaca acaccgagca ccacaaacta gagatc

- 785 <210> SEQ ID NO: 21
- 786 <211> LENGTH: 358
- 787 <212> TYPE: DNA
- 788 <213> ORGANISM: Artificial Sequence
- 790 <220> FEATURE:
- 791 <223> OTHER INFORMATION: Description of Artificial Sequence: MPr1216
- 792 promoter
- 794 <220> FEATURE:
- 795 <221> NAME/KEY: misc feature
- 796 <222> LOCATION: (7)..(14)
- 797 <223> OTHER INFORMATION: G-like box
- 799 <220> FEATURE:

PATENT APPLICATION: US/09/870,375A

DATE: 09/17/2004 TIME: 09:55:41

Input Set : A:\final seq listing.TXT
Output Set: N:\CRF4\09172004\1870375A.raw

```
800 <221> NAME/KEY: enhancer
     801 <222> LOCATION: (39)..(76)
     802 <223> OTHER INFORMATION: Enhancer box
     804 <220> FEATURE:
     805 <221> NAME/KEY: misc feature
     806 <222> LOCATION: (102)..(109)
     807 <223> OTHER INFORMATION: G-like box
     809 <220> FEATURE:
     810 <221> NAME/KEY: enhancer
     811 <222> LOCATION: (134)..(171)
     812 <223> OTHER INFORMATION: Enhancer box
     814 <220> FEATURE:
     815 <221> NAME/KEY: TATA signal
     816 <222> LOCATION: (290)..(296)
     817 <223> OTHER INFORMATION: TATA box
     819 <220> FEATURE:
     820 <221> NAME/KEY: misc feature
     821 <222> LOCATION: (320)
     822 <223> OTHER INFORMATION: Transcription Initiation Site
     824 <400> SEQUENCE: 21
     825 egeogattae gtggetttag cagactgtee aaaaatetgt tittgeaaage tecaattget 60
     826 ccttgcttat ccagcttctt ttgtgttggt ctagacgccg attacgtggc tttagcagac 120
     827 tgtccaaaaa tctgttttgc aaagctccaa ttgctccttg cttatccagc ttcttttgtg 180
     828 ttggcaaact gcgcttttcc aaccgatttt gttcttctcg cgctttcttc ttaggctaaa 240
     829 caaacctcac cgtgcacgca gccatggtcc tgaaccttca cctcgtccct ataaaagcct 300
     830 agccaacctt cacaatctta tcatcaccca caacaccgag caccacaaac tagagatc
E--> 832 <210> SEQ ID NO:
E--> 832 <211> LENGTH:
                              Compet p.4
E--> 832 <212> TYPE:
E--> 832 <213> ORGANISM:
E--> 832 <400> SEQUENCE: 22
E--> 832 22
     833 <211> LENGTH: 453
     834 <212> TYPE: DNA
     835 <213> ORGANISM: Artificial Sequence
W--> 837 <220> FEATURE:
W--> 838 <223> OTHER INFORMATION: Description of Artificial Sequence: MPr1217
     839
               promoter
W--> 841 <220> FEATURE:
W--> 842 <221> NAME/KEY: misc feature
W--> 843 <222> LOCATION: (8)..(15)
W--> 844 <223> OTHER INFORMATION: G-like box
W--> 846 <220> FEATURE:
W--> 847 <221> NAME/KEY: enhancer
W--> 848 <222> LOCATION: (40)..(77)
W--> 849 <223> OTHER INFORMATION: Enhancer box
W--> 851 <220> FEATURE:
W--> 852 <221> NAME/KEY: misc feature
W--> 853 <222> LOCATION: (102)..(109)
```

PATENT APPLICATION: US/09/870,375A

DATE: 09/17/2004 TIME: 09:55:41

Input Set : A:\final seq listing.TXT
Output Set: N:\CRF4\09172004\1870375A.raw

W--> 854 <223> OTHER INFORMATION: G-like box W--> 856 <220> FEATURE: W--> 857 <221> NAME/KEY: enhancer W--> 858 <222> LOCATION: (134)..(171) W--> 859 <223> OTHER INFORMATION: Enhancer box W--> 861 <220> FEATURE: W--> 862 <221> NAME/KEY: misc feature W--> 863 <222> LOCATION: (197)..(204) W--> 864 <223> OTHER INFORMATION: G-like box W--> 866 <220> FEATURE: W--> 867 <221> NAME/KEY: enhancer W--> 868 <222> LOCATION: (229)..(266) W--> 869 <223> OTHER INFORMATION: Enhancer box W--> 871 <220> FEATURE: W--> 872 <221> NAME/KEY: TATA signal W--> 873 <222> LOCATION: (385)..(391) W--> 874 <223> OTHER INFORMATION: TATA box W--> 876 <220> FEATURE: W--> 877 <221> NAME/KEY: misc feature W--> 878 < 222 > LOCATION: (415)W--> 879 <223> OTHER INFORMATION: Transcription Initiation Site E--> 881 <210> SEQ ID NO: E--> 881 <400> SEQUENCE: 22 882 acgccgatta cgtggcttta gcagactgtc caaaaatctg ttttgcaaag ctccaattgc 60 883 teettgetta teeagettet tttgtgttgg tetagaegeg attacgtgge tttageagae 120 884 tgtccaaaaa tctgttttgc aaagctccaa ttgctccttg cttatccagc ttcttttgtg 180 885 ttggtétaga egeegattae gtggetttag cagaetgtee aaaaatetgt tttgcaaage 240 886 tecaattget cettgettat ecagetfett ftgtgttgge aaactgeget tttecaaceg 300 887 attttgttet tetegegett tettettagg etaaacaaac eteacegtge aegeagecat 360 888 ggtcctgaac éttcaccteg tecetataaa ageetageea acettcacaa tettateate 420 889 acccacaaca ccgagcacca caaactagag atc E--> 891 <210> SEQ ID NO: 23 892 <211> LENGTH: 83 893 <212> TYPE: DNA 894 <213> ORGANISM: Artificial Sequence 896 <220> FEATURE: 897 <223> OTHER INFORMATION: Description of Artificial Sequence:Oligodesoxynucleotide 900 <400> SEQUENCE: 23 901 tacqaattcc tcgacatggt tagaagtttt gagtgccgcc actactcgac atggttagaa 60

902 gttttgagtg gccgtagatt tgc

charge to

(<400>)22

<211> 453

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:MPr1217
 promoter

<220>

<221> misc\_feature

<222> (8)..(15)

<223> G-like box

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/17/2004
PATENT APPLICATION: US/09/870,375A TIME: 09:55:42

Input Set : A:\final seq listing.TXT
Output Set: N:\CRF4\09172004\1870375A.raw

#### Skipped Sequences (NEW RULES):

Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

Seq#:22

#### VERIFICATION SUMMARY

DATE: 09/17/2004 PATENT APPLICATION: US/09/870,375A TIME: 09:55:42

Input Set : A:\final seq listing.TXT Output Set: N:\CRF4\09172004\1870375A.raw

```
L:608 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:96 SEQ:16
L:832 M:282 E: Numeric Field Identifier Missing, <210> is required.
L:832 M:282 E: Numeric Field Identifier Missing, <211> is required.
L:832 M:282 E: Numeric Field Identifier Missing, <212> is required.
L:832 M:282 E: Numeric Field Identifier Missing, <213> is required.
L:832 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:21 differs:22
L:832 M:301 E: (44) No Sequence Data was Shown, SEQ ID:0
L:837 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:838 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:841 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:842 M:281 W: Numeric Fields not Ordered, <221> not ordered!.
L:843 M:281 W: Numeric Fields not Ordered, <222> not ordered!.
L:844 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:846 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:847 M:281 W: Numeric Fields not Ordered, <221> not ordered!.
L:848 M:281 W: Numeric Fields not Ordered, <222> not ordered!.
L:849 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:851 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:852 M:281 W: Numeric Fields not Ordered, <221> not ordered!.
L:853 M:281 W: Numeric Fields not Ordered, <222> not ordered!.
L:854 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:856 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:857 M:281 W: Numeric Fields not Ordered, <221> not ordered!.
L:858 M:281 W: Numeric Fields not Ordered, <222> not ordered!.
L:859 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:861 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:862 M:281 W: Numeric Fields not Ordered, <221> not ordered!.
L:863 M:281 W: Numeric Fields not Ordered, <222> not ordered!.
L:864 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:866 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:867 M:281 W: Numeric Fields not Ordered, <221> not ordered!.
L:868 M:281 W: Numeric Fields not Ordered, <222> not ordered!.
L:869 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:871 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:872 M:281 W: Numeric Fields not Ordered, <221> not ordered!.
L:873 M:281 W: Numeric Fields not Ordered, <222> not ordered!.
L:874 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:876 M:281 W: Numeric Fields not Ordered, <220> not ordered!.
L:877 M:281 W: Numeric Fields not Ordered, <221> not ordered!.
L:878 M:281 W: Numeric Fields not Ordered, <222> not ordered!.
L:879 M:281 W: Numeric Fields not Ordered, <223> not ordered!.
L:881 M:282 E: Numeric Field Identifier Missing, <210> is required.
L:881 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:21 differs:22
L:891 M:214 E: (33) Seq.# missing, SEQ ID NO:22
L:1100 M:112 C: (48) String data converted to lower case,
L:1102 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:15 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (38) Counted (39)
```